

AMENDMENTS TO THE SPECIFICATION

Please enter the following amendments to the Specification.

[At page 8, line 17]:

In general, variants will be at least 75% homologous, more suitably at least 80%, preferably at least 85%, and more preferably at least 90% homologous to a polypeptide as for example shown in SEQ ID NO: 2, 4 or 8.

Variant peptides or polypeptides, resulting from rational or established methods of mutagenesis or from combinatorial chemistries as hereinafter described, may comprise conservative amino acid substitutions. Exemplary conservative substitutions in an immuno-interactive polypeptide or polypeptide fragment according to the invention may be made according to the following table:

| TABLE A | |
|------------------|-------------------------|
| Original Residue | Exemplary Substitutions |
| Ala | Ser |
| Arg | Lys |
| Asn | Gln, His |
| Asp | Glu |
| Cys | Ser |
| Gln | Asn |
| Glu | Asp |
| Gly | Pro |
| His | Asn, Gln |
| Ile | Leu, Val |
| Leu | Ile, Val |
| Lys | Arg, Gln, Glu |
| Met | Leu, Ile, |
| Phe | Met, Leu, Tyr |
| Ser | Thr |
| Thr | Ser |
| Trp | Tyr |
| Tyr | Trp, Phe |
| Val | Ile, Leu |

[Following page 28]:

Please Insert the Sequence Listing (pages 1-9)